#2

OIPE

RAW SEQUENCE LISTING DATE: 10/18/2001 PATENT APPLICATION: US/09/965,640 TIME: 09:50:57

Input Set : A:\0315-C Seq Listce filed 092701.txt

Output Set: N:\CRF3\10182001\1965640.raw

	3	<110)> A	PPLI	CANT	: Si	ms,	John	Ε.								I =	INTERI
	5	<120)> T	ITLE	OF	INVE	OITN	N: I	L-1 1	DELT	A DN	A AN	D PO	LYPE	PTIDI	ES		
	7	<130)> F	ILE 1	REFE	RENC	E: 0	315-0	C									
C>						PPLI						/965	,640					
	10	<141	l> CI	URRE	NT F	ILIN	G DA'	re:	2001	-09-	27							
	12	<150)> PI	RIOR	APP:	LICA:	TION	NUM	BER:	09/	612,	921						
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,	15	<160)> и	UMBE	R OF	SEQ	ID I	NOS:	4									
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		caa	aca		nat	gcc	aαt	cta		cct	ata	atc	cta		αtt	caa	ααα	192
						Ala												172
	45	9	50	БСи			DCI	55	DCI	110			60	01	,	01	011	
		σσα		caσ	tac	cta	tet		aaa	aca	σασ	aaa		сса	att	cta	aaa	240
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			gag	cca	ata	aac	atc	atq	gag	ctc	tac	ctc	aaa.	qcc	aaq	qaa	tca	288
						Asn												
	53					85					90					95		
!	55	aag	agc	ttc	acc	ttc	tac	cqq	cqq	gat	atq	ggt	ctt	acc	tcc	agc	ttc	336
						Phe												
	57						_	_	_	_		_			110			
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(60	Glu	Ser	Ala	Ala	Tyr	Pro	Gly	Trp	Phe	Leu	Cys	Thr	Ser	Pro	Glu	Ala	
(51			115					120					125				
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			Gln			Arg												•
(65		130					135					140					
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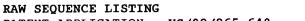
RAW SEQUENCE LISTING

DATE: 10/18/2001 TIME: 09:50:57

PATENT APPLICATION: US/09/965,640

Input Set : A:\0315-C Seq Listce filed 092701.txt
Output Set: N:\CRF3\10182001\I965640.raw

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74 <212> TYPE: PRT	
75 <213> ORGANISM: Mus musculus	
77 <400> SEQUENCE: 2	
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80 1 5 10 15	
83 Leu Lys Val Leu Tyr Leu His Asn Asn Gln Leu Leu Ala Gly Gly Leu	
84 20 25 30	
87 His Ala Glu Lys Val Ile Lys Gly Glu Glu Ile Ser Val Val Pro Asn	
88 35 40 45 91 Arg Ala Leu Asp Ala Ser Leu Ser Pro Val Ile Leu Gly Val Gln Gly	
92 50 55 60 95 Gly Ser Gln Cys Leu Ser Cys Gly Thr Glu Lys Gly Pro Ile Leu Lys	
96 65 70 75 80	
99 Leu Glu Pro Val Asn Ile Met Glu Leu Tyr Leu Gly Ala Lys Glu Ser	
100 85 90 95	
103 Lys Ser Phe Thr Phe Tyr Arg Arg Asp Met Gly Leu Thr Ser Ser Phe	
104 100 105 110	
107 Glu Ser Ala Ala Tyr Pro Gly Trp Phe Leu Cys Thr Ser Pro Glu Ala	
108 115 120 125	
111 Asp Gln Pro Val Arg Leu Thr Gln Ile Pro Glu Asp Pro Ala Trp Asp	
112 130 135 140	
115 Ala Pro Ile Thr Asp Phe Tyr Phe Gln Gln Cys Asp	
116 145 150 155	
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120 <211> LENGTH: 468	
121 <212> TYPE: DNA 122 <213> ORGANISM: Homo sapiens	
122 <213> ORGANISM: HOMO SAPIENS 124 <220> FEATURE:	
124 <220> FEATORE. 125 <221> NAME/KEY: CDS	
126 <222> LOCATION: (1)(468)	
127 <223> OTHER INFORMATION:	
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132 Met Val Leu Ser Gly Ala Leu Cys Phe Arg Met Lys Asp Ser Ala Leu	
133 1 5 10 15	0.6
135 aag gtg ctt tat ctg cat aat aac cag ctt cta gct gga ggg ctg cat	. 96
136 Lys Val Leu Tyr Leu His Asn Asn Gln Leu Leu Ala Gly Gly Leu His	
137 20 25 30	1 / /
139 gca ggg aag gtc att aaa ggt gaa gag atc agc gtg gtc ccc aat cgg	144
140 Ala Gly Lys Val Ile Lys Gly Glu Glu Ile Ser Val Val Pro Asn Arg	
147 22	192
143 tgg ctg gat gcc agc ctg tcc ccc gtc atc ctg ggt gtc cag ggt gga 144 Trp Leu Asp Ala Ser Leu Ser Pro Val Ile Leu Gly Val Gln Gly Gly	170
145 50 55 60 147 agc cag tgc ctg tca tgt ggg gtg ggg cag gag ccg act cta aca cta	240
148 Ser Gln Cys Leu Ser Cys Gly Val Gly Gln Glu Pro Thr Leu Thr Leu	
140 Det off old men ber old off the ord the first the	



PATENT APPLICATION: US/09/965,640 TIME: 09:50:57

DATE: 10/18/2001

Input Set : A:\0315-C Seq Listce filed 092701.txt
Output Set: N:\CRF3\10182001\1965640.raw

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149						70			+-+	a++		~~~	224	~~	+ 00		28	88
151	gag	cca	gtg	aac	atc	atg	gag	CLC	Lat	Tou	21	315	T	Clu	Cor	Luc	2 (00
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153					85					90	a+ a		+00	200		a a a	3:	36
155	agc	ttc	acc	TTC	tac	cgg	cgg	gac	atg	999	CLC	mbm	Con	ayc	Dha	Clu		30
	Ser	Pne	Thr		Tyr	Arg	Arg	Asp	мет 105	СТА	ren	THE	ser	110	Pile	GIU		
157				100			.			+~~	200	~+~	aat		~~	ant.	33	84
159	tcg	gct	gcc	tac	ccg	ggc	Egg	Dha	ctg	Crra	acy mh∞	y L y	Dro	Clu	31a	yaı Acn	3(0 4
	Ser	Ата		туг	Pro	СТУ	ттр		теп	Cys	THI	vaı	125	GIU	на	кър		
161			115					120		~~~	+	aat		+ ~ ~	22+	acc	1	32
	cag																4.	<i>J</i>
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165		130				+	135	~~~	~~~	+~+	~~~						16	68
	CCC											Lay					4.	00
	Pro	тте	Thr	ASP	Pne		Pne	GIII	GTII	Cys	155							
	145	\. a:	70 TI			150					122							
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	•					Homo sapiens												
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179	Met				Gly	Ala	Leu	Cys	Phe		Met	Lys	Asp	Ser		Leu		
179 180	Met 1	Val	Leu	Ser	Gly 5					10					15			
179 180 183	Met	Val	Leu	Ser Tyr	Gly 5				Gln	10				Gly	15			
179 180 183 184	Met 1 Lys	Val Val	Leu Leu	Ser Tyr 20	Gly 5 Leu	His	Asn	Asn	Gln 25	10 Leu	Leu	Ala	Gly	Gly 30	15 Leu	His		
179 180 183 184 187	Met 1	Val Val	Leu Leu Lys	Ser Tyr 20	Gly 5 Leu	His	Asn	Asn Glu	Gln 25	10 Leu	Leu	Ala	Gly v al	Gly 30	15 Leu	His		
179 180 183 184 187	Met 1 Lys Ala	Val Val Gly	Leu Leu Lys 35	Ser Tyr 20 Val	Gly 5 Leu Ile	His Lys	Asn Gly	Asn Glu 40	Gln 25 Glu	10 Leu Ile	Leu Ser	Ala Val	Gly Val 45	Gly 30 Pro	15 Leu Asn	His Arg		
179 180 183 184 187 188 191	Met 1 Lys	Val Val Gly Leu	Leu Leu Lys 35	Ser Tyr 20 Val	Gly 5 Leu Ile	His Lys	Asn Gly Ser	Asn Glu 40	Gln 25 Glu	10 Leu Ile	Leu Ser	Ala Val Gly	Gly Val 45	Gly 30 Pro	15 Leu Asn	His Arg		
179 180 183 184 187 188 191 192	Met 1 Lys Ala Trp	Val Val Gly Leu 50	Leu Lys 35 Asp	Ser Tyr 20 Val Ala	Gly 5 Leu Ile Ser	His Lys Leu	Asn Gly Ser 55	Asn Glu 40 Pro	Gln 25 Glu Val	10 Leu Ile Ile	Leu Ser Leu	Ala Val Gly 60	Gly Val 45 Val	Gly 30 Pro	15 Leu Asn Gly	His Arg Gly		
179 180 183 184 187 188 191 192 195	Met 1 Lys Ala Trp Ser	Val Val Gly Leu 50	Leu Lys 35 Asp	Ser Tyr 20 Val Ala	Gly 5 Leu Ile Ser	His Lys Leu Cys	Asn Gly Ser 55	Asn Glu 40 Pro	Gln 25 Glu Val	10 Leu Ile Ile	Leu Ser Leu Glu	Ala Val Gly 60	Gly Val 45 Val	Gly 30 Pro	15 Leu Asn Gly	His Arg Gly Leu		
179 180 183 184 187 188 191 192 195 196	Met 1 Lys Ala Trp Ser 65	Val Val Gly Leu 50 Gln	Leu Lys 35 Asp Cys	Ser Tyr 20 Val Ala Leu	Gly 5 Leu Ile Ser	His Lys Leu Cys 70	Asn Gly Ser 55 Gly	Asn Glu 40 Pro Val	Gln 25 Glu Val Gly	10 Leu Ile Ile Gln	Leu Ser Leu Glu 75	Ala Val Gly 60 Pro	Gly Val 45 Val Thr	Gly 30 Pro Gln Leu	15 Leu Asn Gly Thr	His Arg Gly Leu 80		
179 180 183 184 187 188 191 192 195 196 199	Met 1 Lys Ala Trp Ser	Val Val Gly Leu 50 Gln	Leu Lys 35 Asp Cys	Ser Tyr 20 Val Ala Leu	Gly 5 Leu Ile Ser Ser	His Lys Leu Cys 70	Asn Gly Ser 55 Gly	Asn Glu 40 Pro Val	Gln 25 Glu Val Gly	10 Leu Ile Ile Gln Leu	Leu Ser Leu Glu 75	Ala Val Gly 60 Pro	Gly Val 45 Val Thr	Gly 30 Pro Gln Leu	15 Leu Asn Gly Thr	His Arg Gly Leu 80		
179 180 183 184 187 188 191 192 195 196 199 200	Met 1 Lys Ala Trp Ser 65 Glu	Val Val Gly Leu 50 Gln Pro	Leu Lys 35 Asp Cys	Ser Tyr 20 Val Ala Leu Asn	Gly 5 Leu Ile Ser Ser Ile 85	His Lys Leu Cys 70 Met	Asn Gly Ser 55 Gly	Asn Glu 40 Pro Val Leu	Gln 25 Glu Val Gly Tyr	10 Leu Ile Ile Gln Leu 90	Leu Ser Leu Glu 75 Gly	Ala Val Gly 60 Pro Ala	Gly Val 45 Val Thr	Gly 30 Pro Gln Leu Glu	15 Leu Asn Gly Thr Ser 95	His Arg Gly Leu 80 Lys		
179 180 183 184 187 188 191 192 195 196 199 200 203	Met 1 Lys Ala Trp Ser 65	Val Val Gly Leu 50 Gln Pro	Leu Lys 35 Asp Cys	Tyr 20 Val Ala Leu Asn	Gly 5 Leu Ile Ser Ser Ile 85	His Lys Leu Cys 70 Met	Asn Gly Ser 55 Gly	Asn Glu 40 Pro Val Leu	Gln 25 Glu Val Gly Tyr Met	10 Leu Ile Ile Gln Leu 90	Leu Ser Leu Glu 75 Gly	Ala Val Gly 60 Pro Ala	Gly Val 45 Val Thr	Gly 30 Pro Gln Leu Glu ser	15 Leu Asn Gly Thr Ser 95	His Arg Gly Leu 80 Lys		
179 180 183 184 187 188 191 192 195 196 199 200 203 204	Met 1 Lys Ala Trp Ser 65 Glu Ser	Val Val Gly Leu 50 Gln Pro	Leu Lys 35 Asp Cys Val Thr	Tyr 20 Val Ala Leu Asn Phe 100	Gly 5 Leu Ile Ser Ser Ile 85 Tyr	His Lys Leu Cys 70 Met	Asn Gly Ser 55 Gly Glu Arg	Asn Glu 40 Pro Val Leu Asp	Gln 25 Glu Val Gly Tyr Met 105	10 Leu Ile Ile Gln Leu 90 Gly	Leu Ser Leu Glu 75 Gly Leu	Ala Val Gly 60 Pro Ala Thr	Gly Val 45 Val Thr Lys Ser	Gly 30 Pro Gln Leu Glu Ser 110	15 Leu Asn Gly Thr Ser 95 Phe	His Arg Gly Leu 80 Lys Glu		
179 180 183 184 187 188 191 192 195 196 199 200 203 204 207	Met 1 Lys Ala Trp Ser 65 Glu	Val Val Gly Leu 50 Gln Pro	Leu Lys 35 Asp Cys Val Thr	Tyr 20 Val Ala Leu Asn Phe 100	Gly 5 Leu Ile Ser Ser Ile 85 Tyr	His Lys Leu Cys 70 Met	Asn Gly Ser 55 Gly Glu Arg	Asn Glu 40 Pro Val Leu Asp	Gln 25 Glu Val Gly Tyr Met 105	10 Leu Ile Ile Gln Leu 90 Gly	Leu Ser Leu Glu 75 Gly Leu	Ala Val Gly 60 Pro Ala Thr	Gly Val 45 Val Thr Lys Ser Pro	Gly 30 Pro Gln Leu Glu Ser 110	15 Leu Asn Gly Thr Ser 95 Phe	His Arg Gly Leu 80 Lys Glu		
179 180 183 184 187 188 191 192 195 196 199 200 203 204 207 208	Met 1 Lys Ala Trp Ser 65 Glu Ser	Val Val Gly Leu 50 Gln Pro Phe Ala	Leu Lys 35 Asp Cys Val Thr Ala 115	Tyr 20 Val Ala Leu Asn Phe 100 Tyr	Gly 5 Leu Ile Ser Ser Ile 85 Tyr Pro	His Lys Leu Cys 70 Met Arg	Asn Gly Ser 55 Gly Glu Arg Trp	Asn Glu 40 Pro Val Leu Asp Phe 120	Gln 25 Glu Val Gly Tyr Met 105 Leu	10 Leu Ile Ile Gln Leu 90 Gly Cys	Leu Ser Leu Glu 75 Gly Leu Thr	Ala Val Gly 60 Pro Ala Thr	Gly Val 45 Val Thr Lys Ser Pro 125	Gly 30 Pro Gln Leu Glu Ser 110 Glu	15 Leu Asn Gly Thr Ser 95 Phe	His Arg Gly Leu 80 Lys Glu Asp		
179 180 183 184 187 188 191 192 195 196 199 200 203 204 207 208 211	Met 1 Lys Ala Trp Ser 65 Glu Ser	Val Val Gly Leu 50 Gln Pro Phe Ala Pro	Leu Lys 35 Asp Cys Val Thr Ala 115	Tyr 20 Val Ala Leu Asn Phe 100 Tyr	Gly 5 Leu Ile Ser Ser Ile 85 Tyr Pro	His Lys Leu Cys 70 Met Arg	Asn Gly Ser 55 Gly Glu Arg Trp Gln	Asn Glu 40 Pro Val Leu Asp Phe 120	Gln 25 Glu Val Gly Tyr Met 105 Leu	10 Leu Ile Ile Gln Leu 90 Gly Cys	Leu Ser Leu Glu 75 Gly Leu Thr	Ala Val Gly 60 Pro Ala Thr Val Gly	Gly Val 45 Val Thr Lys Ser Pro 125	Gly 30 Pro Gln Leu Glu Ser 110 Glu	15 Leu Asn Gly Thr Ser 95 Phe	His Arg Gly Leu 80 Lys Glu Asp		
179 180 183 184 187 188 191 192 195 196 199 200 203 204 207 208 211 212	Met 1 Lys Ala Trp Ser 65 Glu Ser Ser	Val Val Gly Leu 50 Gln Pro Phe Ala Pro 130	Leu Lys 35 Asp Cys Val Thr Ala 115 Val	Tyr 20 Val Ala Leu Asn Phe 100 Tyr	Gly 5 Leu Ile Ser Ser Ile 85 Tyr Pro Leu	His Lys Leu Cys 70 Met Arg Gly Thr	Asn Gly Ser 55 Gly Glu Arg Trp Gln 135	Asn Glu 40 Pro Val Leu Asp Phe 120 Leu	Gln 25 Glu Val Gly Tyr Met 105 Leu	10 Leu Ile Ile Gln Leu 90 Gly Cys	Leu Ser Leu Glu 75 Gly Leu Thr	Ala Val Gly 60 Pro Ala Thr	Gly Val 45 Val Thr Lys Ser Pro 125	Gly 30 Pro Gln Leu Glu Ser 110 Glu	15 Leu Asn Gly Thr Ser 95 Phe	His Arg Gly Leu 80 Lys Glu Asp		
179 180 183 184 187 188 191 192 195 196 199 200 203 204 207 208 211 212 215	Met 1 Lys Ala Trp Ser 65 Glu Ser	Val Val Gly Leu 50 Gln Pro Phe Ala Pro 130	Leu Lys 35 Asp Cys Val Thr Ala 115 Val	Tyr 20 Val Ala Leu Asn Phe 100 Tyr	Gly 5 Leu Ile Ser Ser Ile 85 Tyr Pro Leu	His Lys Leu Cys 70 Met Arg Gly Thr	Asn Gly Ser 55 Gly Glu Arg Trp Gln 135	Asn Glu 40 Pro Val Leu Asp Phe 120 Leu	Gln 25 Glu Val Gly Tyr Met 105 Leu	10 Leu Ile Ile Gln Leu 90 Gly Cys	Leu Ser Leu Glu 75 Gly Leu Thr	Ala Val Gly 60 Pro Ala Thr Val Gly	Gly Val 45 Val Thr Lys Ser Pro 125	Gly 30 Pro Gln Leu Glu Ser 110 Glu	15 Leu Asn Gly Thr Ser 95 Phe	His Arg Gly Leu 80 Lys Glu Asp		

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/965,640

DATE: 10/18/2001 TIME: 09:50:58

Input Set : A:\0315-C Seq Listce filed 092701.txt
Output Set: N:\CRF3\10182001\I965640.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application Number